OIPE

RAW SEQUENCE LISTING   DATE: 11/13/2000   **ENTERED**
PATENT APPLICATION: US/09/698,705   TIME: 12:17:45

     Input Set : A:\P1777R1.txt
     Output Set: N:\CRF3\11132000\I698705.raw

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  3  <110> APPLICANT: Devaux, B.
  4        Keller, G.
  5        Koeppen, H.
  6        Lasky, L.
  8  <120> TITLE OF INVENTION: Anti-Tumor Antibody Compositions and Methods of Use
 10  <130> FILE REFERENCE: P1777R1
W--> 12 <140> CURRENT APPLICATION NUMBER: US/09/698,705
 12  <141> CURRENT FILING DATE: 2000-10-27
 14  <150> PRIOR APPLICATION NUMBER: US 60/162,558
 15  <151> PRIOR FILING DATE: 1999-10-29
 17  <150> PRIOR APPLICATION NUMBER: US 60/182,872
 18  <151> PRIOR FILING DATE: 2000-02-16
 20  <160> NUMBER OF SEQ ID NOS: 25
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 23  <211> LENGTH: 123
 24  <212> TYPE: PRT
 25  <213> ORGANISM: Homo sapiens
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 28  Met Lys Ala Val Leu Leu Ala Leu Leu Met Ala Gly Leu Ala Leu
 29   1           5                   10                      15
 31  Gln Pro Gly Thr Ala Leu Leu Cys Tyr Ser Cys Lys Ala Gln Val
 32                  20                  25                      30
 34  Ser Asn Glu Asp Cys Leu Gln Val Glu Asn Cys Thr Gln Leu Gly
 35                  35                  40                      45
 37  Glu Gln Cys Trp Thr Ala Arg Ile Arg Ala Val Gly Leu Leu Thr
 38                  50                  55                      60
 40  Val Ile Ser Lys Gly Cys Ser Leu Asn Cys Val Asp Asp Ser Gln
 41                  65                  70                      75
 43  Asp Tyr Tyr Val Gly Lys Lys Asn Ile Thr Cys Cys Asp Thr Asp
 44                  80                  85                      90
 46  Leu Cys Asn Ala Ser Gly Ala His Ala Leu Gln Pro Ala Ala Ala
 47                  95                  100                     105
 49  Ile Leu Ala Leu Leu Pro Ala Leu Gly Leu Leu Leu Trp Gly Pro
 50                  110                 115                     120
 52  Gly Gln Leu
 55  <210> SEQ ID NO: 2
 56  <211> LENGTH: 372
 57  <212> TYPE: DNA
 58  <213> ORGANISM: Homo sapiens
 60  <400> SEQUENCE: 2
 61      atg aag gct gtg ctg ctt gcc ctg ttg atg gca ggc 36
 62      Met Lys Ala Val Leu Leu Ala Leu Leu Met Ala Gly
 63       1           5                   10
 65  ttg gcc ctg cag cca ggc act gcc ctg ctg tgc tac tcc 75
 66  Leu Ala Leu Gln Pro Gly Thr Ala Leu Leu Cys Tyr Ser
 67          15                  20                  25
 69  tgc aaa gcc cag gtg agc aac gag gac tgc ctg cag gtg 114
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      70  Cys Lys Ala Gln Val Ser Asn Glu Asp Cys Leu Gln Val
      71                   30                       35
      73  gag aac tgc acc cag ctg ggg gag cag tgc tgg acc gcg 153
      74  Glu Asn Cys Thr Gln Leu Gly Glu Gln Cys Trp Thr Ala
      75          40                  45                  50
      77  cgc atc cgc gca gtt ggc ctc ctg acc gtc atc agc aaa 192
      78  Arg Ile Arg Ala Val Gly Leu Leu Thr Val Ile Ser Lys
      79              55                  60
      81  ggc tgc agc ttg aac tgc gtg gat gac tca cag gac tac 231
      82  Gly Cys Ser Leu Asn Cys Val Asp Asp Ser Gln Asp Tyr
      83    65                  70                   75
      85  tac gtg ggc aag aag aac atc acg tgc tgt gac acc gac 270
      86  Tyr Val Gly Lys Lys Asn Ile Thr Cys Cys Asp Thr Asp
      87              80                  85                  90
      89  ttg tgc aac gcc agc ggg gcc cat gcc ctg cag ccg gct 309
      90  Leu Cys Asn Ala Ser Gly Ala His Ala Leu Gln Pro Ala
      91                   95                  100
      93  gcc gcc atc ctt gcg ctg ctc cct gca ctc ggc ctg ctg 348
      94  Ala Ala Ile Leu Ala Leu Leu Pro Ala Leu Gly Leu Leu
      95          105                 110                 115
      97  ctc tgg gga ccc ggc cag cta tag 372
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      99              120                 124
     101 <210> SEQ ID NO: 3
     102 <211> LENGTH: 96
     103 <212> TYPE: PRT
     104 <213> ORGANISM: Mus musculus
     106 <400> SEQUENCE: 3
     107 Ser Val Ser Ile Ser Cys Arg Ser Ser Lys Ser Leu Leu His Ser
     108  1               5                  10                  15
     110 Asn Gly Asn Thr Tyr Leu Tyr Trp Phe Leu Gln Arg Pro Gly Gln
     111              20                  25                  30
     113 Ser Pro Gln Leu Leu Ile Tyr Arg Met Ser Asn Leu Ala Ser Gly
     114              35                  40                  45
     116 Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Val Phe Thr
     117              50                  55                  60
     119 Leu Arg Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val Tyr Tyr
     120              65                  70                  75
     122 Cys Met Gln His Leu Glu Ser Pro Phe Thr Phe Gly Ser Gly Thr
     123              80                  85                  90
     125 Lys Leu Glu Ile Lys Arg
     126              95
     128 <210> SEQ ID NO: 4
     129 <211> LENGTH: 115
     130 <212> TYPE: PRT
     131 <213> ORGANISM: Mus Musculus
     133 <220> FEATURE:
     134 <221> NAME/KEY: unsure
     135 <222> LOCATION: 46-48, 50-52
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 136 <223> OTHER INFORMATION: unknown amino acid
 138 <400> SEQUENCE: 4
 139   Glu Leu Val Lys Pro Gly Ala Pro Val Lys Leu Ser Cys Lys Ala
 140    1               5                  10                  15
 142   Ser Gly Tyr Thr Phe Thr Asn Tyr Trp Met Asn Trp Val Lys Gln
 143                   20                  25                  30
 145   Arg Pro Gly Arg Gly Leu Glu Trp Ile Gly Arg Ile Asp Pro Ser
 146                   35                  40                  45
W--> 148   Xaa Xaa Xaa Thr Xaa Xaa Xaa Gln Thr Phe Lys Asp Lys Ala Thr
 149                   50                  55                  60
 151   Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr Ile Gln Leu Ser
 152                   65                  70                  75
 154   Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys Ala Ile Thr
 155                   80                  85                  90
 157   Ala Ala Ile Ala Met Asp Tyr Trp Gly Gln Gly Thr Ser Val Thr
 158                   95                 100                 105
 160   Val Ser Ser Ala Lys Thr Thr Gly Pro Ser
 161                  110                 115
 163 <210> SEQ ID NO: 5
 164 <211> LENGTH: 113
 165 <212> TYPE: PRT
 166 <213> ORGANISM: Mus musculus
 168 <400> SEQUENCE: 5
 169   Asp Ile Val Met Thr Gln Ala Ala Pro Ser Val Pro Val Thr Pro
 170    1               5                  10                  15
 172   Gly Glu Ser Val Ser Ile Ser Cys Arg Ser Ser Lys Ser Leu Leu
 173                   20                  25                  30
 175   His Ser Asn Gly Asn Thr Tyr Leu Tyr Trp Phe Leu Gln Arg Pro
 176                   35                  40                  45
 178   Gly Gln Ser Pro Gln Leu Leu Ile Tyr Arg Met Ser Asn Leu Ala
 179                   50                  55                  60
 181   Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly Ser Gly Thr Ala
 182                   65                  70                  75
 184   Phe Thr Leu Arg Ile Ser Arg Val Glu Ala Glu Asp Val Gly Val
 185                   80                  85                  90
 187   Tyr Tyr Cys Leu Gln His Leu Glu Tyr Pro Tyr Thr Phe Gly Gly
 188                   95                 100                 105
 190   Gly Thr Lys Leu Glu Leu Lys Arg
 191                  110
 193 <210> SEQ ID NO: 6
 194 <211> LENGTH: 124
 195 <212> TYPE: PRT
 196 <213> ORGANISM: Mus musculus
 198 <400> SEQUENCE: 6
 199   Gln Val Gln Val Gln Gln Pro Gly Ala Glu Leu Val Lys Pro Gly
 200    1               5                  10                  15
 202   Ala Pro Val Lys Leu Ser Cys Lys Ala Ser Gly Tyr Thr Phe Thr
 203                   20                  25                  30
 205   Asn Tyr Trp Leu Asn Trp Val Lys Gln Arg Pro Gly Arg Gly Leu
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206                    35                  40                  45
208  Glu Trp Ile Gly Arg Ile Asp Pro Ser Asp Ser Glu Ile His Tyr
209                    50                  55                  60
211  Asp Gln Lys Phe Lys Asp Lys Ala Thr Leu Thr Val Asp Lys Ser
212                    65                  70                  75
214  Ser Ser Thr Ala Tyr Ile Gln Leu Ser Ser Leu Thr Ser Glu Asp
215                    80                  85                  90
217  Ser Ala Val Tyr Tyr Cys Ala Leu Thr Gly Ile Tyr Ala Met Ala
218                    95                 100                 105
220  Tyr Trp Gly Gln Gly Thr Ser Val Thr Val Ser Ser Ala Lys Thr
221                   110                 115                 120
223  Thr Gly Pro Ser
226  <210> SEQ ID NO: 7
227  <211> LENGTH: 113
228  <212> TYPE: PRT
229  <213> ORGANISM: Mus musculus
231  <400> SEQUENCE: 7
232  Asp Val Val Met Thr Gln Thr Pro Leu Thr Leu Ser Val Thr Ile
233   1                 5                  10                  15
235  Gly Gln Pro Ala Ser Ile Ser Cys Lys Ser Ser Gln Ser Leu Leu
236                    20                  25                  30
238  Asp Ser Asp Gly Lys Thr Tyr Leu Asn Trp Leu Leu Gln Arg Pro
239                    35                  40                  45
241  Gly Gln Ser Pro Lys Arg Leu Ile Tyr Leu Val Ser Thr Leu Asp
242                    50                  55                  60
244  Ser Gly Val Pro Asp Arg Phe Thr Gly Ser Gly Ser Gly Thr Asp
245                    65                  70                  75
247  Phe Thr Leu Lys Ile Ser Arg Val Glu Ala Glu Asp Leu Gly Val
248                    80                  85                  90
250  Tyr Tyr Cys Trp Gln Gly Thr His Phe Pro Arg Thr Phe Gly Gly
251                    95                 100                 105
253  Gly Thr Lys Leu Glu Ile Lys Arg
254                   110
256  <210> SEQ ID NO: 8
257  <211> LENGTH: 121
258  <212> TYPE: PRT
259  <213> ORGANISM: Mus musculus
261  <400> SEQUENCE: 8
262  Glu Val Gln Leu Gln Gln Ser Gly Pro Asp Leu Glu Lys Pro Gly
263   1                 5                  10                  15
265  Ala Ser Val Lys Ile Ser Cys Lys Pro Ser Gly Asn Ser Phe Thr
266                    20                  25                  30
268  Gly Tyr Tyr Ile His Trp Val Lys Gln Ser His Gly Lys Ser Leu
269                    35                  40                  45
271  Glu Trp Ile Gly Arg Val Asp Pro Asn Asn Gly Phe Thr Ser Tyr
272                    50                  55                  60
274  Asn Gln Lys Phe Lys Gly Lys Ala Ile Leu Thr Val Asp Lys Ser
275                    65                  70                  75
277  Ser Ser Thr Ala Tyr Met Glu Leu Arg Ser Leu Thr Ser Glu Asp
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                    RAW SEQUENCE LISTING              DATE: 11/13/2000
                    PATENT APPLICATION:  US/09/698,705   TIME: 12:17:45

                    Input Set : A:\P1777R1.txt
                    Output Set: N:\CRF3\11132000\I698705.raw

278                     80                   85                   90
280   Ser Ala Val Tyr Tyr Cys Val Gly Asn Phe Phe Asp Ser Trp Gly
281                     95                  100                  105
283   Gln Gly Thr Thr Leu Thr Val Ser Ser Ala Lys Thr Thr Gly Pro
284                    110                  115                  120
286   Ser
289   <210> SEQ ID NO: 9
290   <211> LENGTH: 118
291   <212> TYPE: PRT
292   <213> ORGANISM: Mus musculus
294   <400> SEQUENCE: 9
295   Pro Gly Ala Glu Leu Val Lys Pro Gly Ala Pro Val Lys Leu Ser
296    1                5                   10                   15
298   Cys Lys Ala Ser Gly Tyr Thr Phe Thr Asn Tyr Trp Met Asn Trp
299                     20                   25                   30
301   Val Lys Gln Arg Pro Gly Arg Gly Leu Glu Trp Ile Gly Arg Ile
302                     35                   40                   45
304   Asp Pro Ser Asp Ser Glu Thr Gln Tyr Asn Gln Thr Phe Lys Asp
305                     50                   55                   60
307   Lys Ala Thr Leu Thr Val Asp Lys Ser Ser Ser Thr Ala Tyr Ile
308                     65                   70                   75
310   Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser Ala Val Tyr Tyr Cys
311                     80                   85                   90
313   Ala Ile Thr Ala Ala Ile Ala Met Asp Tyr Trp Gly Gln Gly Thr
314                     95                  100                  105
316   Ser Val Thr Val Ser Ser Ala Lys Thr Thr Gly Pro Ser
317                    110                  115
319   <210> SEQ ID NO: 10
320   <211> LENGTH: 238
321   <212> TYPE: PRT
322   <213> ORGANISM: Artificial sequence
324   <220> FEATURE:
325   <223> OTHER INFORMATION: sequence is chimeric mouse/human
327   <400> SEQUENCE: 10
328   Met Gly Trp Ser Cys Ile Ile Leu Phe Leu Val Ala Thr Ala Thr
329    1                5                   10                   15
331   Gly Val His Ser Asp Ile Val Met Thr Gln Ala Ala Pro Ser Val
332                     20                   25                   30
334   Pro Val Thr Pro Gly Glu Ser Val Ser Ile Ser Cys Arg Ser Ser
335                     35                   40                   45
337   Lys Ser Leu Leu His Ser Asn Gly Asn Thr Tyr Leu Tyr Trp Phe
338                     50                   55                   60
340   Leu Gln Arg Pro Gly Gln Ser Pro Gln Leu Leu Ile Tyr Arg Met
341                     65                   70                   75
343   Ser Asn Leu Ala Ser Gly Val Pro Asp Arg Phe Ser Gly Ser Gly
344                     80                   85                   90
346   Ser Gly Thr Ala Phe Thr Leu Arg Ile Ser Arg Val Glu Ala Glu
347                     95                  100                  105
349   Asp Val Gly Val Tyr Tyr Cys Leu Gln His Leu Glu Tyr Pro Tyr
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VERIFICATION SUMMARY   DATE: 11/13/2000
PATENT APPLICATION: US/09/698,705   TIME: 12:17:46

Input Set : A:\P1777R1.txt
Output Set: N:\CRF3\11132000\I698705.raw

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L:98 M:258 W: Mandatory Feature missing, <220> not found for SEQ ID#:2
L:98 M:258 W: Mandatory Feature missing, <221> not found for SEQ ID#:2
L:98 M:258 W: Mandatory Feature missing, <222> not found for SEQ ID#:2
L:98 M:258 W: Mandatory Feature missing, <223> not found for SEQ ID#:2
L:98 M:340 W: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:2
L:148 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4